

CK330264/c

DEFINITION H8202E07-3 NIA Mouse Unique Gene Set Version 2 Mus musculus cDNA clone H8202E07 3', mRNA sequence.

VERSION CK330264.1 GI:40229789

SOURCE Mus musculus (house mouse)

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia;
Sciurognathi; Muroidea; Muridae; Murinae; Mus.

AUTHORS VanBuren,V., Piao,Y., Dudekula,D.B., Qian,Y., Carter,M.G.,
Martin,P.R., Stagg,C.A., Bassey,U., Aiba,K., Hamatani,T.,
Kargul,G.J., Luo,A.G., Kelso,J., Hide,W. and Ko,M.S.H.

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Plate: H8202 row: E column: 07

Seq primer: -21M13 Forward

High quality sequence stop: 465

POLYA=Yes.

source

1. .465

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/organism="Mus musculus"
```

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/mol_type="mRNA"
```

```
/strain="C57BL/6"
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/db_xref="niaEST:H8202E07-3"
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/db xref="taxon:10090"
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/clone="H8202E07"
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/sex="mixed"
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```
/dev stage="mixed"
```

```
/lab_host="DH10B"
```

```
/clone lib="NIA Mouse Unique Gene Set Version 2"
```

```
/note="Vector: pSPORT1 (Invitrogen); Site_1: SalI; Site_2:
```

NotI; Mouse cDNA project by the Laboratory of Genetics,

National Institute on Aging (NIA), Intramural Research

Program, NIH (<http://lgsun.grc.nia.nih.gov/cDNA>). This

clone is among a rearranged set of 11,424 clones from more

than 20 cDNA libraries."

Query Match 98.9%; Score 86; DB 8; Length 465;

Best Local Similarity 100.0%; Pred. No. 8.4e-18;

Matches 86; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 GCAGCCCCTGGGACCCCTCCGATAGATCTGCCCTCTTGAAAAGCAAAGCTGAGGGCTCTGC 61

[illegible]

Db 446 GCAGCCCCTGGGACCCCTCCGATAGATCTGCCCTCTTGAAAAGCAAACCTGAGGGGCTCTGC 387

Qy	62	TCGCTGGCCCTCGGAGCCTACGAAGA	87
Db	386	TCGCTGGCCCTCGGAGCCTACGAAGA	361